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Claims

What is claimed is:

1	1.	A method for categorizing a dataset comprising a plurality of datapoints, each
2		datapoint comprising at least two numerical values, said method comprising the
3		steps of:
4		(a) producing a plurality of angular values by calculating an angular value for
5		each datapoint based on said datapoint's numerical values;
6		(b) sorting said plurality of datapoints by said angular values;
7		(c) producing a plurality of difference values by calculating differences
8		between adjacent angular values;
9		(d) determining at least one category-dividing value by identifying at least one
10		difference value above a predetermined threshold gap value; and
11		(e) classifying at least one datapoint according to its angular value relative to
12		at least one category-dividing value.
1	2.	The method of claim 1 wherein each datapoint comprises two numerical values.
1	3.	The method of claim 2 wherein said angular value is an arctangent of said two
2		numerical values.
1	4.	The method of claim 1 wherein said numerical values represent fluorometric data.
1	5.	The method of claim 1 wherein said determining step (d) identifies two category-
2		dividing values.
1	6.	The method of claim 1 further comprising the step of normalizing said numerical
2		values to a scale.

The method of claim 6 wherein said scale ranges from 0.0 to 1.0.

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- 1 8. The method of claim 1 further comprising the step of removing non-amplification datapoints from said dataset, said step comprising the steps of:
 - (i) calculating a Euclidean distance for each datapoint;
- removing at least one datapoint from said dataset, wherein the Euclidean distance of said datapoint falls below a predetermined distance threshold.
- The method of claim 1 wherein said determining step (d) identifies two category-dividing values comprising a first and a second category-dividing value, and said classifying step (e) comprises the steps of:
 - classifying at least one datapoint in a first category, wherein all datapoints
 of said first category have an angular value lower than said first and
 second category-dividing values;
 - (ii) classifying at least one datapoint in a second category, wherein all datapoints of said second category have an angular value between said first and second category-dividing values; and
 - (iii) classifying at least one datapoint in a third category, wherein all datapoints of said third category have an angular value greater than said first and second category-dividing values.
- 1 10. The method of claim 9 wherein classification in said first category corresponds to
 2 homozygosity for a first allele, classification in said third category corresponds to
 3 homozygosity for a second allele, and classification in said second category
 4 corresponds to heterozygosity for said first and second alleles.
- 1 11. The method of claim 10 further comprising the step of determining the presence 2 of a condition to bring to the attention of a human user, wherein said condition 3 comprises the proportion of datapoints classified as heterozygous exceeding a 4 predetermined threshold.

- 1 12. The method of claim 11 further comprising the step of determining the presence of a condition to bring to the attention of a human user.
- 1 13. The method of claim 12 wherein said condition comprises a substantial majority of datapoints being classified in one category.
- 1 14. The method of claim 13 wherein said category corresponds to heterozygosity for a first and second allele.
- 1 15. The method of claim 13 wherein said category corresponds to homozygosity for either a first or second allele.
- 1 16. The method of claim 13 wherein said category cannot be determined to correspond to either heterozygosity or homozygosity.
- 1 17. The method of claim 12 wherein said condition comprises said datapoints being classified into more than three categories.
- 1 18. The method of claim 12 wherein said condition comprises at least one of said datapoints remaining unclassified.
- 1 19. The method of claim 12 wherein said condition comprises the Euclidean distance 2 between at least one of said classified datapoints and at least one non-3 amplification datapoint being below a predetermined threshold.
- The method of claim 12 wherein said condition comprises a substantial majority of datapoints in said first category having an angular value higher than a predetermined threshold.
- 1 21. The method of claim 20 wherein said angular value is an arctangent and said predetermined threshold is 0.67.

- The method of claim 12 wherein said condition comprises a substantial majority of datapoints in said third category having an angular value lower than a predetermined threshold.
- The method of claim 22 wherein said angular value is an arctangent and said predetermined threshold is 1.0.
- The method of claim 12 wherein said condition comprises a substantial majority of datapoints in said second category having an angular value lower than a first predetermined threshold or higher than a second predetermined threshold.
- The method of claim 24 wherein said angular value is an arctangent, said first predetermined threshold is 0.18, and said second predetermined threshold is 1.35.
- The method of claim 12 wherein said condition comprises the difference between the largest angular value of a datapoint in a category and the smallest angular value of a datapoint in the category exceeding a predetermined threshold.
- The method of claim 26 wherein said angular value is an arctangent and said second predetermined threshold is 0.6.
- The method of claim 12 wherein said first allele is a major allele and said second allele is a minor allele, and said major and minor alleles are in a Hardy-Weinberg equilibrium.
- The method of claim 28 further comprising the step of determining the presence of a condition to bring to the attention of a human user, wherein said condition indicates an incompatibility with a Hardy-Weinberg equilibrium.

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- The method of claim 29 wherein said incompatibility comprises a greater number of datapoints classified as homozygous for said minor allele than classified as heterozygous.
- The method of claim 12 further comprising the step of determining the presence of a condition to bring to the attention of a human user, said determining step comprising the steps of:
- calculating the center of the set of removed datapoints, said center comprising an x and y coordinate; and
 - (ii) determining if either said x or y coordinate exceeds a predetermined threshold.
 - 32. The method of claim 31 wherein said predetermined threshold is 0.3 on a normalized scale of 0.0 to 1.0.